PF-0576 USN

```
<110> HILLMAN, Jennifer L.; YUE, Henry
    CORLEY, Neil C.; GUEGLER, Karl J.
    PATTERSON, Chandra
```

- <120> EXTRACELLULAR ADHESIVE PROTEINS, EXADH1 AND EXADH2
- <130> PF-0576 USN
- <140> US 09/762,527
- <141> To Be Assigned
- <150> PCT/US99/17997
- <151> 1999-08-09
- <150> US 09/131,648
- <151> 1998-08-10
- <160> 5
- <170> PERL Program
- <210> 1
- <211> 336
- <212> PRT
- <213> Homo sapiens
- <220>
- <221> misc\_feature
- <223> Incyte ID No: 2635136
- <400> 1
- Met Ser Gln Pro Ser Gly Gly Arg Ala Pro Gly Thr Arg Ile Tyr
  1 5 10 15
- Ser Trp Ser Cys Pro Thr Val Met Ser Pro Gly Glu Lys Leu Asp
- Pro Ile Pro Asp Ser Phe Ile Leu Gln Pro Pro Val Phe His Pro 35 40 45
- Val Val Pro Tyr Val Thr Thr Ile Phe Gly Gly Leu His Ala Gly
  50 55 60
- Lys Met Val Met Leu Gln Gly Val Val Pro Leu Asp Ala His Arg
- 65 70 75
  Phe Gln Val Asp Phe Gln Cys Gly Cys Ser Leu Cys Pro Arg Pro
- 80 85 90 Asp Ile Ala Phe His Phe Asn Pro Arg Phe His Thr Thr Lys Pro
- 95 100 105
  His Val Ile Cys Asn Thr Leu His Gly Gly Arg Trp Gln Arg Glu
- Ala Arg Trp Pro His Leu Ala Leu Arg Arg Gly Ser Ser Phe Leu
- $125 \hspace{1cm} 130 \hspace{1cm} 135$  Ile Leu Phe Leu Phe Gly Asn Glu Glu Val Lys Val Ser Val Asn
- 140 145 150 Gly Gln His Phe Leu His Phe Arg Tyr Arg Leu Pro Leu Ser His
- 155 160 165
- Val Asp Thr Leu Gly Ile Phe Gly Asp Ile Leu Val Glu Ala Val 170 175 180
- Gly Phe Leu Asn Ile Asn Pro Phe Val Glu Gly Ser Arg Glu Tyr 185 190 195
- Pro Ala Gly His Pro Phe Leu Leu Met Ser Pro Arg Leu Glu Val 200 205 210
- Pro Cys Ser His Ala Leu Pro Gln Gly Leu Ser Pro Gly Gln Val

```
215
                                    220
Ile Ile Val Arg Gly Leu Val Leu Gln Glu Pro Lys His Phe Thr
Val Ser Leu Arg Asp Gln Ala Ala His Ala Pro Val Thr Leu Arg
                245
                                    250
Ala Ser Phe Ala Asp Arg Thr Leu Ala Trp Ile Ser Arg Trp Gly
                260
                                    265
Gln Lys Lys Leu Ile Ser Ala Pro Phe Leu Phe Tyr Pro Gln Arg
                275
                                    280
Phe Phe Glu Val Leu Leu Phe Gln Glu Gly Gly Leu Lys Leu
                290
                                     295
Ala Leu Asn Gly Gln Gly Leu Gly Ala Thr Ser Met Asn Gln Gln
                305
                                    310
Ala Leu Glu Gln Leu Arg Glu Leu Arg Ile Ser Gly Ser Val Gln
                                    325
Leu Tyr Cys Val His Ser
<210> 2
<211> 708
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: 2687731
<400> 2
Met Lys Asp Met Pro Leu Arg Ile His Val Leu Leu Gly Leu Ala
Ile Thr Thr Leu Val Gln Ala Val Asp Lys Lys Val Asp Cys Pro
Arg Leu Cys Thr Cys Glu Ile Arg Pro Trp Phe Thr Pro Arg Ser
Ile Tyr Met Glu Ala Ser Thr Val Asp Cys Asn Asp Leu Gly Leu
                 50
Leu Thr Phe Pro Ala Arg Leu Pro Ala Asn Thr Gln Ile Leu Leu
                                     70
Leu Gln Thr Asn Asn Ile Ala Lys Ile Glu Tyr Ser Thr Asp Phe
                 80
                                     85
Pro Val Asn Leu Thr Gly Leu Asp Leu Ser Gln Asn Asn Leu Ser
                 95
                                    100
Ser Val Thr Asn Ile Asn Val Lys Lys Met Pro Gln Leu Leu Ser
                110
                                    115
Val Tyr Leu Glu Glu Asn Lys Leu Thr Glu Leu Pro Glu Lys Cys
                                    130
Leu Ser Glu Leu Ser Asn Leu Gln Glu Leu Tyr Ile Asn His Asn
Leu Leu Ser Thr Ile Ser Pro Gly Ala Phe Ile Gly Leu His Asn
                155
                                    160
Leu Leu Arg Leu His Leu Asn Ser Asn Arg Leu Gln Met Ile Asn
                170
                                    175
Ser Lys Trp Phe Asp Ala Leu Pro Asn Leu Glu Ile Leu Met Ile
                185
                                    190
Gly Glu Asn Pro Ile Ile Arg Ile Lys Asp Met Asn Phe Lys Pro
                200
                                    205
Leu Ile Asn Leu Arg Ser Leu Val Ile Ala Gly Ile Asn Leu Thr
                215
                                    220
Glu Ile Pro Asp Asn Ala Leu Val Gly Leu Glu Asn Leu Glu Ser
```

				230					235					240
Ile	Ser	Phe	Tyr	Asp 245	Asn	Arg	Leu	Ile	Lys 250	Val	Pro	His	Val	Ala 255
Leu	Gln	Lys	Val	Val 260	Asn	Leu	Lys	Phe	Leu 265	Asp	Ľeu	Asn	Lys	Asn 270
Pro	Ile	Asn	Arg	Ile 275	Arg	Arg	Gly	Asp	Phe 280	Ser	Asn	Met	Leu	His 285
Leu	Lys	Glu	Leu	Gly 290	Ile	Asn	Asn	Met	Pro 295	Glu	Leu	Ile	Ser	Ile 300
Asp	Ser	Leu	Ala	Val 305	Asp	Asn	Leu	Pro	Asp 310	Leu	Arg	Lys	Ile	Glu 315
Ala	Thr	Asn	Asn	Pro 320	Arg	Leu	Ser	Tyr	Ile 325	His	Pro	Asn	Ala	Phe 330
Phe	Arg	Leu	Pro	Lys 335	Leu	Glu	Ser	Leu	Met 340	Leu	Asn	Ser	Asn	Ala 345
Leu	Ser	Ala	Leu	Tyr 350	His	Gly	Thr	Ile	Glu 355	Ser	Leu	Pro	Asn	Leu 360
Lys	Glu	Ile	Ser	Ile 365	His	Ser	Asn	Pro	Ile 370	Arg	Cys	Asp	Cys	Val 375
Ile	Arg	Trp	Met	Asn 380	Met	Asn	Lys	Thr	Asn 385	Ile	Arg	Phe	Met	Glu 390
Pro	Asp	Ser	Leu	Phe 395	Суѕ	Val	Asp	Pro	Pro 400	Glu	Phe	Gln	Gly	Gln 405
Asn	Val	Arg	Gln	Val 410	His	Phe	Arg	Asp	Met 415	Met	Glu	Ile	Сув	Leu 420
Pro	Leu	Ile	Ala	Pro 425	Glu	Ser	Phe	Pro	Ser 430	Asn	Leu	Asn	Val	Glu 435
Ala	Gly	Ser	Tyr	Val 440	Ser	Phe	His	Cys	Arg 445	Ala	Thr	Ala	Glu	Pro 450
Gln	Pro	Glu	Ile	Туr 455	Trp	Ile	Thr	Pro	Ser 460	Gly	Gln	Lys	Leu	Leu 465
Pro	Asn	Thr	Leu	Thr 470	Asp	Lys	Phe	Tyr	Val 475	His	Ser	Glu	Gly	Thr 480
Leu	Asp	Ile	Asn	Gly 485	Val	Thr	Pro	Lys	Glu 490	Gly	Gly	Leu	Tyr	Thr 495
				500					505				Val	510
				515					520				Ser	525
				530					535				Val	540
				545					550				Trp	555
				560					565				Ala	570
				575					580				Asn	585
				590					595				Tyr	600
				605					610				Leu	615
				620					625				Leu	630
				635	,				640				Сув	645
				650					655				His	660
Tyr	Val	Arg	Asn	Tyr 665	Leu	Gln	Lys	Pro	Thr 670	Phe	Ala	Leu	Gly	Glu 675

```
Leu Tyr Pro Pro Leu Ile Asn Leu Trp Glu Ala Gly Lys Glu Lys
                                    685
Ser Thr Ser Leu Lys Val Lys Ala Thr Val Ile Gly Leu Pro Thr
                695
                                    700
                                                        705
Asn Met Ser
<210> 3
<211> 1643
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: 2635136
<400> 3
tgcaatggcc atatgctgca gacccggagt gggtagttag ttggttaatg ccagtcttcc 60
tcccctggac actgagttct gctgacagcc cccgcccagc cagagctctg ctgtatacca 120
ccgggagtgg ggctggtgt gagcctggag gtcgcccgct gccctcctag ggctgctcca 180
gacagcatta aaacgctgca ggtcgcaggt gagactaaca gctgggagag ctgctccagg 240
catttaggac cctgactggg gcagatgagt cagcccagtg ggggcagggc tcctggaacg 300
aggatctaca gttggagttg ccccactgtc atgtcacctg gagaaaaact ggacccaatt 360
cctgacagct tcattctgca accaccagtc ttccacccgg tggttcctta tgtcacgacg 420
atttttggag gcctgcatgc aggcaagatg gtcatgctgc aaggagtggt ccctctagat 480
gcacacaggt ttcaggtgga cttccagtgt ggctgcagcc tgtgtccccg gccagatatc 540
geettecact teaacceteg ettecatace accaageece atgteatetg caacaccetg 600
catggtggac gctggcaaag ggaggcccgg tggccccacc tggccctgcg aagaggctcc 660
agetteetea teetetttet ettegggaat gaggaagtga aggtgagtgt gaatggacag 720
cactttetee actteegeta eeggeteeca etgteteatg tggacaeget gggtatattt 780
ggtgacatcc tggtagaggc tgttggattc ctgaacatca atccatttgt ggagggcagc 840
agagagtace cagetggaca teettteetg etgatgagee eeaggetgga ggtgeeetge 900
tcacatgctc ttccccaggg tctctcgcct gggcaggtca tcatagtacg gggactggtc 960
ttgcaagagc cgaagcattt tactgtgagc ctgagggacc aggctgccca tgctcctgtg 1020
acactcaggg ceteettege agacagaact etggeetgga tetecegetg ggggeagaag 1080
aaactgatct cagccccctt cctcttttac ccccagagat tctttgaggt gctgctcctg 1140
ttccaggagg gagggctgaa gctggcgctc aatgggcagg ggctgggggc caccagcatg 1200
aaccagcagg ccctggagca gctgcgggag ctccggatca gtggaagtgt ccagctctac 1260
tgtgtccact cctgaggatg gttccaggga aataccgcca gaaaacaaga aggtcagccc 1320
acteceaggg ecceaetete eteceeteat taaaceatee acetgacace ageacateag 1380
gcctggttca cctctggggt cacgagactg agtctacagg agctttgggc ctgagggaag 1440
gcacaagagt gcaaaggttc ctcgaactct gcaccttcct ccaccaggag cctgggatat 1500
ggctccatct gccttcaggg cctggactgc actcacagag gcaagtgttg tagactaaca 1560
aagatactcc aaaatacaat ggcttaaaga atgtggtcat ttattcttta ttatttattt 1620
atttgtggtc aaataaataa ata
                                                                  1643
<210> 4
<211> 2290
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte ID No: 2687731
<400> 4
cttactagca ctgactgtgg aatccttaag ggcccattac atttctgaag aagaaagcta 60
agatgaagga catgccactc cgaattcatg tgctacttgg cctagctatc actacactag 120
tacaagctgt agataaaaaa gtggattgtc cacggttatg tacgtgtgaa atcaggcctt 180
```

```
ggtttacacc cagatccatt tatatggaag catctacagt ggattgtaat gatttaggtc 240
ttttaacttt cccagccaga ttgccagcta acacacagat tcttctccta cagactaaca 300
atattgcaaa aattgaatac tccacagact ttccagtaaa ccttactggc ctggatttat 360
ctcaaaacaa tttatcttca gtcaccaata ttaatgtaaa aaagatgcct cagctccttt 420
ctgtgtacct agaggaaaac aaacttactg aactgcctga aaaatgtctg tccgaactga 480
gcaacttaca agaactctat attaatcaca acttgctttc tacaatttca cctggaqcct 540
ttattggcct acataatctt cttcgacttc atctcaattc aaatagattg cagatgatca 600
acagtaagtg gtttgatgct cttccaaatc tagagattct gatgattggg gaaaatccaa 660
ttatcagaat caaagacatg aactttaagc ctcttatcaa tcttcgcagc ctggttatag 720
ctggtataaa cctcacagaa ataccagata acgccttggt tggactggaa aacttagaaa 780
gcatctcttt ttacgataac aggcttatta aagtacccca tgttgctctt caaaaagttg 840
taaatctcaa atttttggat ctaaataaaa atcctattaa tagaatacga aggggtgatt 900
ttagcaatat gctacactta aaagagttgg ggataaataa tatgcctgag ctgatttcca 960
tcgatagtct tgctgtggat aacctgccag atttaagaaa aatagaagct actaacaacc 1020
ctagattgtc ttacattcac cccaatgcat ttttcagact ccccaagctg gaatcactca 1080
tgctgaacag caatgctctc agtgccctgt accatggtac cattgagtct ctgccaaacc 1140
tcaaggaaat cagcatacac agtaacccca tcaggtgtga ctgtgtcatc cgttggatga 1200
acatgaacaa aaccaacatt cgattcatgg agccagattc actgttttgc gtggacccac 1260
ctgaattcca aggtcagaat gttcggcaag tgcatttcag ggacatgatg gaaatttgtc 1320
tecetettat ageteetgag agettteett etaatetaaa tgtagaaget gggagetatg 1380
tttcctttca ctgtagagct actgcagaac cacagcctga aatctactgg ataacacctt 1440
ctggtcaaaa actcttgcct aataccctga cagacaagtt ctatgtccat tctgagggaa 1500
cactagatat aaatggcgta actcccaaag aagggggttt atatacttgt atagcaacta 1560
acctagttgg cgctgacttg aagtctgtta tgatcaaagt ggatggatct tttccacaag 1620
ataacaatgg ctctttgaat attaaaataa gagatattca ggccaattca gttttggtgt 1680
cctggaaagc aagttctaaa attctcaaat ctagtgttaa atggacagcc tttgtcaaga 1740
ctgaaaattc tcatgctgcg caaagtgctc gaataccatc tgatgtcaag gtatataatc 1800
ttactcatct gaatccatca actgagtata aaatttgtat tgatattccc accatctatc 1860
agaaaaacag aaaaaaatgt gtaaatgtca ccaccaaagg tttgcaccct gatcaaaaag 1920
agtatgaaaa gaataatacc acaacactta tggcctgtct tggaggcctt ctggggatta 1980
ttggtgtgat atgtcttatc agctgcctct ctccagaaat gaactgtgat ggtggacaca 2040
gctatgtgag gaattactta cagaaaccaa cctttgcatt aggtgagctt tatcctcctc 2100
tgataaatct ctgggaagca ggaaaagaaa aaagtacatc actgaaagta aaagcaactg 2160
ttataggttt accaacaat atgtcctaaa aaccaccaag gaaacctact ccaaaaatga 2220
aaaaaaaaa
                                                                 2290
<210> 5
<211> 316
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> GenBank ID No: g1932712
<400> 5
Met Leu Ser Leu Asn Asn Leu Gln Asn Ile Ile Tyr Asn Pro Val
                                    10
                                                        15
Ile Pro Phe Val Gly Thr Ile Pro Asp Gln Leu Asp Pro Gly Thr
                20
                                    25
                                                        30
Leu Ile Val Ile Arg Gly His Val Pro Ser Asp Ala Asp Arg Phe
                35
                                    40
                                                        45
Gln Val Asp Leu Gln Asn Gly Ser Ser Val Lys Pro Arg Ala Asp
                50
Val Ala Phe His Phe Asn Pro Arg Phe Lys Arg Ala Gly Cys Ile
                65
                                                        75
                                    70
Val Cys Asn Thr Leu Ile Asn Glu Lys Trp Gly Arg Glu Glu Ile
```

85

80

## PF-0576 USN

Thr	Tyr	Asp	Thr	Pro 95	Phe	Lys	Arg	Glu	Lys 100	Ser	Phe	Glu	Ile	Val 105
Ile	Met	Val	Leu	Lys 110	Asp	Lys	Phe	Gln		Ala	Val	Asn	Gly	
His	Thr	Leu	Leu		Gly	His	Arg	Ile		Pro	Glu	Lys	Ile	
Thr	Leu	Gly	Ile	Tyr	Gly	Lys	Val	Asn	Ile	His	Ser	Ile	Gly	Phe
Ser	Phe	Ser	Ser	_	Leu	Gln	Ser	Thr		Ala	Ser	Ser	Leu	
Leu	Thr	Glu	Ile		Arg	Glu	Asn	Val		Lys	Ser	Gly	Thr	
Gln	Leu	Ser	Leu		Phe	Ala	Ala	Arg		Asn	Thr	Pro	Met	_
Pro	Gly	Arg	Thr		Val	Val	Gln	Gly		Val	Asn	Ala	Asn	
Lys	Ser	Phe	Asn		Asp	Leu	Leu	Ala	_	Lys	Ser	Lys	Asp	
Ala	Leu	His	Leu		Pro	Arg	Leu	Asn		Lys	Ala	Phe	Val	_
Asn	Ser	Phe	Leu		Glu	Ser	Trp	Gly		Glu	Glu	Arg	Asn	
Thr	Ser	Phe	Pro		Ser	Pro	Gly	Met	-	Phe	Glu	Met	Ile	
Tyr	Cys	Asp	Val	_	Glu	Phe	Lys	Val		Val	Asn	Gly	Val	
Ser	Leu	Glu	Tyr		His	Arg	Phe	Lys		Leu	Ser	Ser	Ile	_
Thr	Leu	Glu	Ile		Gly	Asp	Ile	His		Leu	Glu	Val	Arg	
Trp				305					310					315